OM of: US-08-962-560A-4 to: Issued_Patents_NA:* out_format : pfs

Date: Feb 4, 2000 4:34 PM

About: Results were produced by the GenCore software, version 4.5,
Copyright (c) 1993-2000 Compugen Ltd.

Command line parameters:
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Search information block:
Query: US-08-962-560A-4
Query length: 212
Database: Issued_Patents_NA:*
Database sequences: 207703
Database length: 57918730
Search time (sec): 62.100000

score_list:

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seq_name: /cgn1_6/ptodata/2/ina/5C_COMB.seq:US-08-463-081B-9

seq_documentation_block:
; Sequence 9, Application US/08463081B
; Patent No. 5871960
; Patent No. 5871960 5837487
; GENERAL INFORMATION:
; APPLICANT: Smith, Kendall A. & Beadling, Carol
; TITLE OF INVENTION: Nucleic Acids Encoding CR5 Polypeptide,
; TITLE OF INVENTION: Vector and Transformed Cell Thereof, and Expression Thereof
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: PRETTY, SCHROEDER & POPLAWSKI
; STREET: 444 South Flower St. - Suite 1900
; CITY: Los Angeles
; STATE: California
; COUNTRY: USA
; ZIP: 90071
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0,
; SOFTWARE: Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/463,081B
; FILING DATE: 5-JUN-1995
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/104,736
; FILING DATE: 10-AUG-1993
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/796,066
; FILING DATE: 20-NOV-91
; ATTORNEY/AGENT INFORMATION:
; NAME: Viviana Amzel, Ph. D.
; REGISTRATION NUMBER: 30,930
; REFERENCE/DOCKET NUMBER: P66 38150 (DART-060)
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 622-7700
; TELEFAX: (213) 489-4210
; INFORMATION FOR SEQ ID NO: 9:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1960 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 112..886
US-08-463-081B-9

alignment_scores:
Quality: 249.50 Length: 297
Ratio: 1.835 Gaps: 10
Percent Similarity: 45.791 Percent Identity: 27.946

alignment_block:
US-08-962-560A-4 x US-08-463-081B-9 ..

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; Patent No. 5871961
;  GENERAL INFORMATION:
;    APPLICANT:  Smith, Kendall A. & Beadling, Carol
;    TITLE OF INVENTION:  Nucleic Acids Encoding CR5 Polypeptide,
;    TITLE OF INVENTION:  Vector and Transformed Cell Thereof, and
;    TITLE OF INVENTION:  Expression Thereof
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Ratner & Prestia
;      ADDRESSEE:  .          (B) STREET:One Westlakes-Berwyn
;      CITY:  Valley Forge
;      STATE:  Pennsylvania
;      COUNTRY:  USA
;      ZIP:  19482
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0,
;      SOFTWARE:  Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/461,379A
;      FILING DATE:  5-JUNE-1995
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  USSN 08/330,108; 08/104,736
;      APPLICATION NUMBER:  & 07/796,066
;      FILING DATE:  27-OCT-1994; 10-AUG-1993 & 20-NOV-91
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Viviana Amzel, Ph. D.
;      REGISTRATION NUMBER:  30,930
;      REFERENCE/DOCKET NUMBER:  DART-070
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (610)470-0700
;      TELEFAX:  (610)470-0701
;  INFORMATION FOR SEQ ID NO:  9:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1960 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  112..886
US-08-461-379A-9
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            Quality:  249.50              Length:    297
              Ratio:  1.835                 Gaps:     10
 Percent Similarity:  45.791   Percent Identity: 27.946

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; Sequence 9, Application US/08462390B
; Patent No. 5882894
;  GENERAL INFORMATION:
;    APPLICANT:  Smith, K. A., & Beadling, C.
;    TITLE OF INVENTION:  Nucleic Acids Encoding CR8 Polypeptide, Vector and
;    TITLE OF INVENTION:  Transformed Cell Thereof, and Expression Thereof
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Ratner & Prestia
;      ADDRESSEE:  .          (B) STREET:One Westlakes-Berwyn
;      CITY:  Valley Forge
;      STATE:  Pennsylvania
;      COUNTRY:  USA
;      ZIP:  19482
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/462,390B
;      FILING DATE:  5-JUNE-1995
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  USSN 08/330,108
;      FILING DATE:  27-OCT-1994
;      APPLICATION NUMBER:  USSN 08/104,736
;      FILING DATE:  10-AUG-1993
;      APPLICATION NUMBER:  USSN 07/796,066
;      FILING DATE:  20-NOV-91
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Viviana Amzel, Ph. D.
;      REGISTRATION NUMBER:  30,930
;      REFERENCE/DOCKET NUMBER:  DART-040
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (610)407-0700
;      TELEFAX:  (610)407-0701
;  INFORMATION FOR SEQ ID NO:  9:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1960 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  112..886
US-08-462-390B-9


alignment_scores:
           Quality:  249.50              Length:    297
             Ratio:  1.835                 Gaps:     10
 Percent Similarity:  45.791   Percent Identity: 27.946

alignment_block:
 US-08-962-560A-4 x US-08-462-390B-9    ..

 Align seg 1/1  to: US-08-462-390B-9  from: 1  to: 1960

      14 SerProAlaAlaGluProArgArgArgSerGluProSerSerSerSerSe 30
         :::|||:::|||   |||   |||      :::|||         |||||
      10 GCCCCGGGAGCCTACCCAGCACGCGCTCCGCGCCCACTGGTTCCCTCCAG 59

      30 rSerSerSerProAla...AlaProValArg..................... 39
         |          ||||||   :::|||:::|||
      60 CCGCCGCCGTCCAGCCGAGTCCCCACTCCGGAGTCGCCGCTGCCGCGGGG 109

      40 .......................ProArgProCysProAlaVal...... 46
                                |||||||||      ||||||
     110 ACATGGTCCTCTGCGTTCAGGGACCTCGTCCTTTGCTGGCTGTGGAGCGG 159

      46 ................................................ 46

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      47 .............ProAla.............................. 48
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     210 CATGCAGCCCTTGCCTGCTGGGGCCTTCCTCGAGGAGGTGGCAGAGGGTA 259

      49 ..ProAlaProGlyAspThrHisPheArgThrPheArgSerHisSerAsp 64
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     410 CAGAAGGCACGTTCTTAGTACGTGACAGCACGCACCCCAGCTACCTGTTC 459

     115 AlaLeuSerValLysMetAlaSerGlyProThrSerIleArgValHisPh 131
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     460 ACGCTGTCAGTGAAAACCACTCGTGGCCCCACCAATGTACGCATTGAGTA 509

     131 eGlnAlaGlyArgPheHisLeuAspGly.........SerArgGlu.... 143
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     144 .....ThrPheAspCysLeuPheGluLeuLeuGluHisTyrValAla... 157
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     162 tLeuGlyAlaPro..................................... 166
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     167 ............................................LeuArg 168
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seq_documentation_block:
; Sequence 2, Application US/08918206
; Patent No. 5919661
;  GENERAL INFORMATION:
;    APPLICANT:  Hillman, Jennifer L.
;    APPLICANT:  Guegler, Karl
;    APPLICANT:  Corley, Neil C.
;    APPLICANT:  Shah, Purvi
;    TITLE OF INVENTION:  CYTOKINE INDUCIBLE REGULATORY
;    TITLE OF INVENTION:  PROTEIN
;    NUMBER OF SEQUENCES:  3
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Incyte Pharmaceuticals, Inc.
;      STREET:  3174 Porter Dr.
;      CITY:  Palo Alto
;      STATE:  CA
;      COUNTRY:  USA
;      ZIP:  94304
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette
;      COMPUTER:  IBM Compatible
;      OPERATING SYSTEM:  DOS
;      SOFTWARE:  FastSEQ for Windows Version 2.0
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/918,206
;      FILING DATE:  Filed Herewith
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:
;      FILING DATE:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Billings, Lucy J.
;      REGISTRATION NUMBER:  36,749
;      REFERENCE/DOCKET NUMBER:  PF-0372 US
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415-855-0555
;      TELEFAX:  415-845-4166
;  INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2587 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    IMMEDIATE SOURCE:
;      LIBRARY:  BRSTNOT13
;      CLONE:  2787140
US-08-918-206-2


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            Quality:  240.50              Length:    299
              Ratio:  1.743                 Gaps:     10
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     281 AGCCCCATCCCTTCTCTCTCTGCCACCATTTCGGACACCCCGCAGGAACT 330

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     331 CGTTTTGGGATTCGCACTGACTTCAAGGAAGGACGCGAACCCTTCTCTGA 380

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     481 CCAGTGGGGGACCGCGGGGTCGGCGGAGGAGCCATCCCGCAGGCGGCGCG 530

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     531 TCTGGCGAAGGCTGCGGGAGCTCGGTCAGACAGGATGGTACTGGGGAAGT 580

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     928 GCTCACCATTAACAAATGTACCGGTGCC......ATCTGGGGACTGCCTT 971

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seq_documentation_block:
; Sequence 33, Application US/08463081B
; Patent No. 5871960
; Patent No. 5871960 5837487
;  GENERAL INFORMATION:
;    APPLICANT:  Smith, Kendall A. & Beadling, Carol
;    TITLE OF INVENTION:  Nucleic Acids Encoding CR5 Polypeptide,
;    TITLE OF INVENTION:  Vector and Transformed Cell Thereof, and Expression Thereof
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  PRETTY, SCHROEDER & POPLAWSKI
;      STREET:  444 South Flower St. - Suite 1900
;      CITY:  Los Angeles
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  90071
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0,
;      SOFTWARE:  Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/463,081B
;      FILING DATE:  5-JUN-1995
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/104,736
;      FILING DATE:  10-AUG-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/796,066
;      FILING DATE:  20-NOV-91
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Viviana Amzel, Ph. D.
;      REGISTRATION NUMBER:  30,930
;      REFERENCE/DOCKET NUMBER:  P66 38150 (DART-060)
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (213) 622-7700
;      TELEFAX:  (213) 489-4210
;  INFORMATION FOR SEQ ID NO:  33:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  774 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
US-08-463-081B-33


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            Quality:  237.00            Length:    251
              Ratio:  1.837               Gaps:      7
 Percent Similarity:  51.394   Percent Identity: 29.084

alignment_block:
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seq_name: /cgn1_6/ptodata/2/ina/5C_COMB.seq:US-08-461-379A-33

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; Sequence 33, Application US/08461379A
; Patent No. 5871961
;  GENERAL INFORMATION:
;    APPLICANT:  Smith, Kendall A. & Beadling, Carol
;    TITLE OF INVENTION:  Nucleic Acids Encoding CR5 Polypeptide,
;    TITLE OF INVENTION:  Vector and Transformed Cell Thereof, and
;    TITLE OF INVENTION:  Expression Thereof
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Ratner & Prestia
;      ADDRESSEE:  .          (B) STREET:One Westlakes-Berwyn
;      CITY:  Valley Forge
;      STATE:  Pennsylvania
;      COUNTRY:  USA
;      ZIP:  19482
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0,
;      SOFTWARE:  Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/461,379A
;      FILING DATE:  5-JUNE-1995
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  USSN 08/330,108; 08/104,736
;      APPLICATION NUMBER:  & 07/796,066
;      FILING DATE:  27-OCT-1994; 10-AUG-1993 & 20-NOV-91
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Viviana Amzel, Ph. D.
;      REGISTRATION NUMBER:  30,930
;      REFERENCE/DOCKET NUMBER:  DART-070
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (610)470-0700
;      TELEFAX:  (610)470-0701
;  INFORMATION FOR SEQ ID NO:  33:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  774 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
US-08-461-379A-33


alignment_scores:
           Quality:  237.00            Length:    251
             Ratio:  1.837               Gaps:      7
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; Sequence 33, Application US/08462390B
; Patent No. 5882894
;  GENERAL INFORMATION:
;    APPLICANT:  Smith, K. A., & Beadling, C.
;    TITLE OF INVENTION:  Nucleic Acids Encoding CR8 Polypeptide, Vector and
;    TITLE OF INVENTION:  Transformed Cell Thereof, and Expression Thereof
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Ratner & Prestia
;      ADDRESSEE:  .          (B) STREET:One Westlakes-Berwyn
;      CITY:  Valley Forge
;      STATE:  Pennsylvania
;      COUNTRY:  USA
;      ZIP:  19482
;    COMPUTER READABLE FORM:
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;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/462,390B
;      FILING DATE:  5-JUNE-1995
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  USSN 08/330,108
;      FILING DATE:  27-OCT-1994
;      APPLICATION NUMBER:  USSN 08/104,736
;      FILING DATE:  10-AUG-1993
;      APPLICATION NUMBER:  USSN 07/796,066
;      FILING DATE:  20-NOV-91
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Viviana Amzel, Ph. D.
;      REGISTRATION NUMBER:  30,930
;      REFERENCE/DOCKET NUMBER:  DART-040
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (610)407-0700
;      TELEFAX:  (610)407-0701
;  INFORMATION FOR SEQ ID NO:  33:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  774 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
US-08-462-390B-33


alignment_scores:
            Quality:  237.00              Length:    251
              Ratio:  1.837                 Gaps:      7
 Percent Similarity:  51.394   Percent Identity: 29.084

alignment_block:
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 Align seg 1/1  to: US-08-462-390B-33  from: 1  to: 774

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; Sequence 1, Application US/08259264
; Patent No. 5650293
;  GENERAL INFORMATION:
;    APPLICANT:  White, Morris F.
;    TITLE OF INVENTION:  pp60PIK: A DOWNSTREAM ELEMENT IN INSULIN SIGNALING
;    NUMBER OF SEQUENCES:  2
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  LAHIVE & COCKFIELD
;      STREET:  60 STATE STREET, SUITE 510
;      CITY:  BOSTON
;      STATE:  MASSACHUSETTS
;      COUNTRY:  USA
;      ZIP:  02109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  ASCII text
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/259,264
;      FILING DATE:  10-JUN-1994
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Myers, Louis
;      REGISTRATION NUMBER:  35,965
;      REFERENCE/DOCKET NUMBER:  JDP-021
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617) 227-7400
;      TELEFAX:  (617) 227-5941
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  5737 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
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;      LOCATION:  1388..2749
US-08-259-264-1


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              Ratio:  1.392                Gaps:      8
 Percent Similarity:  53.012   Percent Identity: 27.108

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 Align seg 1/1  to: US-08-259-264-1  from: 1  to: 5737

      34 ProAlaAlaProValArgProArgProCysProAlaValProAlaProAl 50
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    1469 CCAGCTCTTCCACCAAAGCCA...CCTAAGCCAATGACTCCAGCAGTCAC 1515

      50 aProGly......AspThrHisPheArgThrPheArgSerHisSerAspT 65
         :   |||      |||:::   :::
    1516 AAATGGAATGAAGGACAGTTTCATT..................... 1540

      65 yrArgArgIleThrArgThrSerAlaLeuLeuAspAlaCysGlyPheTyr 81
                               :::|||   ||||||      :::|||
    1541 .....................TCTCTTCAAGATGCA...GAGTGGTAC 1564

      82 TrpGlyProLeuSerValHisGlyAlaHisGluArgLeuArgAlaGluPr 98
         ||||||   :::|||   :::   ||||||||||||||||||      ||
    1565 TGGGGAGATATTTCCAGGGAAGAGGTAAATGACAAATTGCGGGACATGCC 1614

      98 oValGlyThrPheLeuValArgAsp...SerArgGlnArgAsnCysPheP 114
         |   ||||||||||||||||||||   |||   :::   :::      :
    1615 AGATGGTACCTTCTTAGTTCGTGATGCCTCAACGAAAATGCAGGGGGATT 1664

     114 heAlaLeuSerValLysMetAlaSerGlyProThrSerIleArgValHis 130
         :::::|||:::::::::   :::::::::          |||:::::::::
    1665 ATACATTGACTTTGAGGAAGGGAGGAAATAATAAATTAATAAAGATCTAT 1714

     131 PheGlnAlaGlyArgPheHisLeuAspGlySerArgGlu......ThrPh 145
            :::   |||::::::         |||      |||      |||||
    1715 CATCGGGATGGTAAATAT.........GGCTTCTCTGAGCCCCTGACGTT 1755

     145 eAspCysLeuPheGluLeuLeuGluHisTyrValAlaAlaProArgArgM 162
         |      :::   ||||||::::::||||||                  :
    1756 TACTTCTGTGGTGGAGCTTATTAACCACTACCACCACGAGTCTCTCGCTC 1805

     162 etLeuGlyAlaProLeuArgGlnArgArgValArgProLeu......... 175
         ::   :::      |||      :::   :::   |||:::
    1806 AGTACAATCCCAAACTCGACGTGAAGCTGACGTACCCAGTATCCAGATTC 1855

     176 ......GlnGluLeuCysArgGlnArgIleValAlaAlaValGlyArg 189
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seq_name: /cgn1_6/ptodata/2/ina/5A_COMB.seq:US-08-076-011-1

seq_documentation_block:
; Sequence 1, Application US/08076011
; Patent No. 5521069
;  GENERAL INFORMATION:
;    APPLICANT:  ONDA, Haruo
;    APPLICANT:  KIMURA, Chiharu
;    APPLICANT:  OHKUBO, Shoichi
;    TITLE OF INVENTION:  NOVEL DNA AND USE THEREOF
;    NUMBER OF SEQUENCES:  3
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  DAVID G. CONLIN; DIKE, BRONSTEIN,
;      ADDRESSEE:  ROBERTS & CUSHMAN
;      STREET:  130 Water Street
;      CITY:  Boston
;      STATE:  Massachusetts
;      COUNTRY:  US
;      ZIP:  02109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/076,011
;      FILING DATE:  11-JUN-1993
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/047,246
;      FILING DATE:  13-APR-1993
;      NAME:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/741,676
;      FILING DATE:  07-AUG-1991
;      NAME:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  RESNICK, DAVID S
;      REGISTRATION NUMBER:  34235
;      REFERENCE/DOCKET NUMBER:  41155-CIP
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617)523-3400
;      TELEFAX:  (617)523-6440
;      TELEX:  200291 STRE UR
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  17041 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  join(7540..7650, 9814..9945, 10421..10519,
;      LOCATION:  11602..11787)
US-08-076-011-1


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              Ratio:  1.235                Gaps:     10
 Percent Similarity:  43.946   Percent Identity: 27.803

alignment_block:
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 Align seg 1/1  to: US-08-076-011-1  from: 1  to: 17041

       2 ValAlaArgAsnGlnValAlaAlaAspAsnAlaIleSerProAlaAlaGl 18
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    6696 GTAGCAGCAGCAGAAGCCGCAGCTTCAGAGGCAGCCGGAGAGACCTCGGA 6745

      18 uProArgArgArgSerGluProSerSerSerSerSerSerSerSerProA 35
         |   |||||||||   :::|||                  |||      |||:
    6746 GCAGAGAAGGCGCCGCCGACCC..............TCGCGGCTGCCTG 6780

      35 laAlaProValArgProArgProCysProAlaValProAlaProAlaPro 51
         ::      :::      |||   :::         |||:::|||   |||
    6781 GCCCGCGGCTCCTACAAAGGCGGGCTAGCCGCCCGCCCTCTCCCTTGCCT 6830

      52 GlyAspThrHisPheArgThrPheArgSerHisSerAspTyrArgArgIl 68
         ::::::      |||   ||||||
    6831 TCCTCCCCTTCTTTTCTGACTTTC..................... 6854

      68 eThrArgThrSerAlaLeuLeuAspAlaCysGlyPheTyrTrpGlyProL 85
                                                      ||||
    6855 .....................................CCTC 6858

      85 euSerValHisGlyAlaHisGluArgLeuArgAlaGluProValGly... 100
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    6859 TTTCCCTTAAT...........CGCCTGCTTCTTCCTCCGGGTGGACTT 6896
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     101 ......ThrPheLeuValArgAspSerArgGlnArgAsnCysPhePheAl 115
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    6897 ACGGCCACCTTGCTCCTCCGCGCTTCACCTCATCGCCCCCTCTTT..... 6941

     115 aLeuSerValLysMetAlaSerGlyProThrSerIleArgValHisPheG 132
          ||||||           |||:::|||   |||:::||||||   :::
    6942 .CTTTCTTCTGCCTCTCTCTCTGCGCCCCCTTCTCTCCGTGTCACGCTCC 6990

     132 lnAlaGlyArgPheHisLeuAspGlySerArgGluThrPheAspCysLeu 148
            |||      :::|||           :::||||||:::
    6991 CTCCTGGTTCTGCGCGTCTA.....CAAACTTTTGAGCAGAAC 7028

     149 PheGluLeuGluHisTyrValAlaAlaProArgArgMetLeuGlyAl 165
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    7029 ACGAGCCTCGGCAAACGA.....GTCCCGCAGCTCCTCCTGCTGCT 7069

     165 aProLeu.....A 168
          ||||||
    7070 CCCGCTGGTTCCTGCGGCTTCTGCTCAGACACCAACGCCAGACGGCGATG 7119

     168 rgGlnArgArgValArgProLeuGlnGlu.....LeuCysArg 180
          :::|||   |||   |||   ||||||          ||||||
    7120 CCTCTCGGGTGGTGACTCCAGCGCAGGAACTTGAAGAAGCGCTTTGCCCG 7169

     181 GlnArgIleValAlaAlaVal.....Gl 188
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    7170 CCGTCCTACCTGGCAGCTCTCCTGGCAGCGGGAGGAGTTGAAGGGTAAGG 7219

     188 yArgGluAsnLeuAlaArg 194
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seq_name: /cgn1_6/ptodata/2/ina/5D_COMB.seq:US-08-850-993-1

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seq_documentation_block:
; Sequence 1, Application US/08850993
; Patent No. 5955277
; GENERAL INFORMATION:
;  APPLICANT: Hansen, Torben
;  APPLICANT: Andersen, Carsten
;  APPLICANT: Pedersen, Oluf B.
;  TITLE OF INVENTION: Mutant cDNA Encoding The p85alpha
;  TITLE OF INVENTION: Subunit Of Phosphatidylinositol 3-Kinase
;  FILE REFERENCE: 4802.200-US
;  CURRENT APPLICATION NUMBER: US/08/850,993
;  CURRENT FILING DATE: 1997-05-05
;  EARLIER APPLICATION NUMBER: 0539/96
;  EARLIER FILING DATE: 1996-05-06
;  NUMBER OF SEQ ID NOS: 25
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 1
;   LENGTH: 2508
;   TYPE: DNA
;   ORGANISM: human
US-08-850-993-1


alignment_scores:
            Quality:  119.50            Length:    179
              Ratio:  1.258               Gaps:      7
 Percent Similarity:  53.073   Percent Identity: 24.581

alignment_block:
 US-08-962-560A-4 x US-08-850-993-1   ..

 Align seg 1/1  to: US-08-850-993-1  from: 1  to: 2508

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     985 .ACTGTAGCCAACAACGGT.....ATGAATAACAATATGT 1018

      74 euLeuAspAlaCysGlyPheTyrTrpGlyProLeuSerValHisGlyAla 90
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    1019 CCTTACAAAATGCTGAATGGTACTGGGGAGATATCTCGAGGGAAGAAGTG 1068

      91 HisGluArgLeuArgAlaGluProValGlyThrPheLeuValArgAsp.. 106
         :::|||:::||||||            ||||||||||||||||||||
    1069 AATGAAAAACTTCGAGATACAGCAGACGGGACCTTTTTGGTACGAGATGC 1118

     107 .SerArgGlnArgAsnCysPhePheAlaLeuSerValLysMetAlaSerG 123
          |||   :::   :::      ::::::|||:::::::::   :::::::
    1119 GTCTACTAAAATGCATGGTGATTATACTCTTACACTAAGGAAAGGGGGAA 1168

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    1169 ATAACAAATTAATCAAAATA.....TTTCATCGAGAT 1200

     140 GlySerArgGlu.....ThrPheAspCysLeuPheGluLe 151
         |||:::                 ||||||:::   :::   |||||
    1201 GGGAAATATGGCTTCTCTGACCCATTAACCTTCAGTTCTGTGGTTGAATT 1250

     151 uLeuGluHisTyrValAlaAlaProArgArgMetLeuGlyAlaProLeuA 168
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    1251 AATAAACCACTACCGGAATGAATCTCTAGCTCAGTATAATCCCAAATTGG 1300

     168 rgGlnArgArgValArgProLeuGlnGluLeuCysArgGlnArgIleVal 184
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    1301 ATGTGAAATTACTTTATCCAGTATCCAAATACCAACAGGATCAAGTTGTC 1350

     185 .............AlaAlaValGlyArgGluAsnLeuAlaArgIlePr 196
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    1351 AAAGAAGATAATATTGAAGCTGTAGGGAAAAAA..... 1383

     196 oLeuAsnProValLeuArgAspTyrLeuSerSerPhe 208
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    1384 ............TTACATGAATATAACACTCAGTTT 1407
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seq_name: /cgn1_6/ptodata/2/ina/5A_COMB.seq:US-07-906-349A-1

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seq_documentation_block:
; Sequence 1, Application US/07906349A
; Patent No. 5434064
;  GENERAL INFORMATION:
;    APPLICANT:  Schlessinger, Joseph
;    APPLICANT:  Skolnik, Edward Y.
;    APPLICANT:  Margolis, Benjamin L.
;    TITLE OF INVENTION:  A NOVEL EXPRESSION-CLONING METHOD FOR
;    TITLE OF INVENTION:  IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE KINASES
;    TITLE OF INVENTION:  TARGET PROTEINS
;    NUMBER OF SEQUENCES:  16
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Browdy and Neimark
;      STREET:  419 Seventh Street, N.W.
;      CITY:  Washington
;      STATE:  D.C.
;      COUNTRY:  USA
;      ZIP:  20004
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/906,349A
;      FILING DATE:  30-JUN-1992
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  07/643,237
;      FILING DATE:  18-JAN-1991
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  202-628-5197
;      TELEFAX:  202-737-3528
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;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3372 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-07-906-349A-1


alignment_scores:
            Quality:  119.50            Length:    179
              Ratio:  1.258               Gaps:      7
 Percent Similarity:  53.073   Percent Identity: 24.581

alignment_block:
 US-08-962-560A-4 x US-07-906-349A-1   ..

 Align seg 1/1  to: US-07-906-349A-1  from: 1  to: 3372

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      57 gThrPheArgSerHisSerAspTyrArgArgIleThrArgThrSerAlaL 74
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     988 .ACTGTAGCCAACAACGGT..............ATGAATAACAATATGT 1021

      74 euLeuAspAlaCysGlyPheTyrTrpGlyProLeuSerValHisGlyAla 90
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    1022 CCTTACAAAATGCTGAATGGTACTGGGGAGATATCTCGAGGGAAGAAGTG 1071

      91 HisGluArgLeuArgAlaGluProValGlyThrPheLeuValArgAsp.. 106
         :::|||:::||||||           ||||||||||||||||||||
    1072 AATGAAAAACTTCGAGATACAGCAGACGGGACCTTTTTGGTACGAGATGC 1121

     107 .SerArgGlnArgAsnCysPhePheAlaLeuSerValLysMetAlaSerG 123
          |||   :::   :::      :::::::|||:::::::::   :::::::
    1122 GTCTACTAAAATGCATGGTGATTATACTCTTACACTAAGGAAAGGGGGAA 1171

     123 lyProThrSerIleArgValHisPheGlnAlaGlyArgPheHisLeuAsp 139
         ::        |||::::::                  ||||||   |||
    1172 ATAACAAATTAATCAAAATA.................TTTCATCGAGAT 1203

     140 GlySerArgGlu..............ThrPheAspCysLeuPheGluLe 151
         |||:::                   |||||||:::   :::   |||||
    1204 GGGAAATATGGCTTCTCTGACCCATTAACCTTCAGTTCTGTGGTTGAATT 1253

     151 uLeuGluHisTyrValAlaAlaProArgArgMetLeuGlyAlaProLeuA 168
         |:::::::||||||                  :::   :::      |||
    1254 AATAAACCACTACCGGAATGAATCTCTAGCTCAGTATAATCCCAAATTGG 1303

     168 rgGlnArgArgValArgProLeuGlnGluLeuCysArgGlnArgIleVal 184
           :::   :::   |||:::::::::     :::::::::::::|||
    1304 ATGTGAAATTACTTTATCCAGTATCCAAATACCAACAGGATCAAGTTGTC 1353

     185 ..............AlaAlaValGlyArgGluAsnLeuAlaArgIlePr 196
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    1354 AAAGAAGATAATATTGAAGCTGTAGGGAAAAAA................. 1386

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seq_name: /cgn1_6/ptodata/2/ina/5A_COMB.seq:US-08-167-035-1

seq_documentation_block:
; Sequence 1, Application US/08167035
; Patent No. 5618691
;  GENERAL INFORMATION:
;    APPLICANT:  Schlessinger, Joseph
;    APPLICANT:  Skolnick, Edward Y.
;    APPLICANT:  Margolis, Benjamin L.
;    TITLE OF INVENTION:  NOVEL EXPRESSION CLONING METHOD FOR
;    TITLE OF INVENTION:  IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE
;    TITLE OF INVENTION:  KINASES AND NOVEL TARGET PROTEINS
;    NUMBER OF SEQUENCES:  50
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  PENNIE & EDMONDS
;      STREET:  1155 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  10036-2711
;      ZIP:  10036-2711
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/167,035
;      FILING DATE:  16-DEC-1993
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Coruzzi, Laura A.
;      REGISTRATION NUMBER:  30,742
;      REFERENCE/DOCKET NUMBER:  7683-062
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 790-9090
;      TELEFAX:  (212) 869-9741/8864
;      TELEX:  66141 PENNIE
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3372 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  43..2214
US-08-167-035-1


alignment_scores:
            Quality:  119.50            Length:    179
              Ratio:  1.258               Gaps:      7
 Percent Similarity:  53.073   Percent Identity: 24.581

alignment_block:
 US-08-962-560A-4 x US-08-167-035-1   ..

 Align seg 1/1  to: US-08-167-035-1  from: 1  to: 3372

      41 ArgProCysProAlaValProAlaProAlaProGlyAspThrHisPheAr 57
         :::|||:::||||||:::|||         |||      |||
     946 CAGCCTGCACCAGCACTGCCTCCTAAACCACCAAAACCTACT........ 987

      57 gThrPheArgSerHisSerAspTyrArgArgIleThrArgThrSerAlaL 74
         |||      :::::::::                    :::::::::
     988 .ACTGTAGCCAACAACGGT..............ATGAATAACAATATGT 1021

      74 euLeuAspAlaCysGlyPheTyrTrpGlyProLeuSerValHisGlyAla 90
           |||:::   :::   :::|||||||||   :::|||   :::   :::
    1022 CCTTACAAAATGCTGAATGGTACTGGGGAGATATCTCGAGGGAAGAAGTG 1071

      91 HisGluArgLeuArgAlaGluProValGlyThrPheLeuValArgAsp.. 106
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    1072 AATGAAAAACTTCGAGATACAGCAGACGGGACCTTTTTGGTACGAGATGC 1121

     107 .SerArgGlnArgAsnCysPhePheAlaLeuSerValLysMetAlaSerG 123
          |||   :::   :::      :::::::|||:::::::::   :::::::
    1122 GTCTACTAAAATGCATGGTGATTATACTCTTACACTAAGGAAAGGGGGAA 1171

     123 lyProThrSerIleArgValHisPheGlnAlaGlyArgPheHisLeuAsp 139
         ::        |||::::::                  ||||||   |||
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 1172 ATAACAAATTAATCAAAATA.................TTTCATCGAGAT 1203

  140 GlySerArgGlu..............ThrPheAspCysLeuPheGluLe 151
      |||:::                   |||||||:::   :::   |||||
 1204 GGGAAATATGGCTTCTCTGACCCATTAACCTTCAGTTCTGTGGTTGAATT 1253

  151 uLeuGluHisTyrValAlaAlaProArgArgMetLeuGlyAlaProLeuA 168
      |:::::::||||||                  :::   :::      |||
 1254 AATAAACCACTACCGGAATGAATCTCTAGCTCAGTATAATCCCAAATTGG 1303

  168 rgGlnArgArgValArgProLeuGlnGluLeuCysArgGlnArgIleVal 184
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 1304 ATGTGAAATTACTTTATCCAGTATCCAAATACCAACAGGATCAAGTTGTC 1353

  185 ..............AlaAlaValGlyArgGluAsnLeuAlaArgIlePr 196
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seq_name: /cgn1_6/ptodata/2/ina/5A_COMB.seq:US-08-167-035-48

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seq_documentation_block:
; Sequence 48, Application US/08167035
; Patent No. 5618691
;  GENERAL INFORMATION:
;    APPLICANT:  Schlessinger, Joseph
;    APPLICANT:  Skolnick, Edward Y.
;    APPLICANT:  Margolis, Benjamin L.
;    TITLE OF INVENTION:  NOVEL EXPRESSION CLONING METHOD FOR
;    TITLE OF INVENTION:  IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE
;    TITLE OF INVENTION:  KINASES AND NOVEL TARGET PROTEINS
;    NUMBER OF SEQUENCES:  50
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  PENNIE & EDMONDS
;      STREET:  1155 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  10036-2711
;      ZIP:  10036-2711
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/167,035
;      FILING DATE:  16-DEC-1993
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Coruzzi, Laura A.
;      REGISTRATION NUMBER:  30,742
;      REFERENCE/DOCKET NUMBER:  7683-062
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 790-9090
;      TELEFAX:  (212) 869-9741/8864
;      TELEX:  66141 PENNIE
;  INFORMATION FOR SEQ ID NO:  48:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3372 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
US-08-167-035-48


 alignment_scores:
            Quality:  119.50              Length:    179
              Ratio:  1.258                Gaps:      7
 Percent Similarity:  53.073   Percent Identity:  24.581
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alignment_block:
 US-08-962-560A-4 x US-08-167-035-48/rev   ..

 Align seg 1/1  to reverse of: US-08-167-035-48  from: 1  to: 3372

      41 ArgProCysProAlaValProAlaProAlaProGlyAspThrHisPheAr 57
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      57 gThrPheArgSerHisSerAspTyrArgArgIleThrArgThrSerAlaL 74
          |||      :::::::::                 :::::::::
    2385 .ACTGTAGCCAACAACGGT..............ATGAATAACAATATGT 2352

      74 euLeuAspAlaCysGlyPheTyrTrpGlyProLeuSerValHisGlyAla 90
          |||:::   :::   :::||||||||||   :::|||   :::   :::
    2351 CCTTACAAAATGCTGAATGGTACTGGGGAGATATCTCGAGGGAAGAAGTG 2302

      91 HisGluArgLeuArgAlaGluProValGlyThrPheLeuValArgAsp.. 106
         :::|||:::||||||              ||||||||||||||||||||
    2301 AATGAAAAACTTCGAGATACAGCAGACGGGACCTTTTTGGTACGAGATGC 2252

     107 .SerArgGlnArgAsnCysPhePheAlaLeuSerValLysMetAlaSerG 123
          |||   :::   :::      ::::::|||:::::::::   :::::::
    2251 GTCTACTAAAATGCATGGTGATTATACTCTTACACTAAGGAAAGGGGGAA 2202

     123 lyProThrSerIleArgValHisPheGlnAlaGlyArgPheHisLeuAsp 139
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    2201 ATAACAAATTAATCAAAATA.................TTTCATCGAGAT 2170

     140 GlySerArgGlu..............ThrPheAspCysLeuPheGluLe 151
         |||:::                   |||||||:::   :::   |||||
    2169 GGGAAATATGGCTTCTCTGACCCATTAACCTTCAGTTCTGTGGTTGAATT 2120

     151 uLeuGluHisTyrValAlaAlaProArgArgMetLeuGlyAlaProLeuA 168
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    2119 AATAAACCACTACCGGAATGAATCTCTAGCTCAGTATAATCCCAAATTGG 2070

     168 rgGlnArgArgValArgProLeuGlnGluLeuCysArgGlnArgIleVal 184
            :::   :::   |||:::::::::      :::::::::::::|||
    2069 ATGTGAAATTACTTTATCCAGTATCCAAATACCAACAGGATCAAGTTGTC 2020

     185 ..............AlaAlaValGlyArgGluAsnLeuAlaArgIlePr 196
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    2019 AAAGAAGATAATATTGAAGCTGTAGGGAAAAAA..................  1987

     196 oLeuAsnProValLeuArgAspTyrLeuSerSerPhe 208
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    1986 .............TTACATGAATATAACACTCAGTTT 1963
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seq_name: /cgn1_6/ptodata/2/ina/5A_COMB.seq:US-08-208-887A-1

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seq_documentation_block:
; Sequence 1, Application US/08208887A
; Patent No. 5677421
;  GENERAL INFORMATION:
;    APPLICANT:  Schlessinger, Joseph
;    APPLICANT:  Skolnick, Edward Y.
;    APPLICANT:  Margolis, Benjamin L.
;    TITLE OF INVENTION:  NOVEL EXPRESSION CLONING METHOD FOR
;    TITLE OF INVENTION:  IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE
;    TITLE OF INVENTION:  KINASES AND NOVEL TARGET PROTEINS
;    NUMBER OF SEQUENCES:  51
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  PENNIE & EDMONDS
;      STREET:  1155 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  10036-2711
;      ZIP:  10036-2711
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
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;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  PatentIn Release #1.0, Version #1.30
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/208,887A
;       FILING DATE:  11-MAR-1994
;       CLASSIFICATION:  435
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Coruzzi, Laura A.
;       REGISTRATION NUMBER:  30,742
;       REFERENCE/DOCKET NUMBER:  7683-063
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  (212) 790-9090
;       TELEFAX:  (212) 869-9741/8864
;       TELEX:  66141 PENNIE
;   INFORMATION FOR SEQ ID NO:  1:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  3372 base pairs
;       TYPE:  nucleic acid
;       STRANDEDNESS:  double
;       TOPOLOGY:  unknown
;     MOLECULE TYPE:  cDNA
;     FEATURE:
;       NAME/KEY:  CDS
;       LOCATION:  43..2214
US-08-208-887A-1


alignment_scores:
            Quality:  119.50                Length:    179
              Ratio:  1.258                   Gaps:      7
 Percent Similarity:  53.073    Percent Identity: 24.581

alignment_block:
 US-08-962-560A-4 x US-08-208-887A-1    ..

 Align seg 1/1  to: US-08-208-887A-1  from: 1  to: 3372

      41 ArgProCysProAlaValProAlaProAlaProGlyAspThrHisPheAr 57
         :::|||:::|||||||:::|||          |||      |||
     946 CAGCCTGCACCAGCACTGCCTCCTAAACCACCAAAACCTACT..... 987

      57 gThrPheArgSerHisSerAspTyrArgArgIleThrArgThrSerAlaL 74
          |||      :::::::::                 :::::::::
     988 .ACTGTAGCCAACAACGGT.............ATGAATAACAATATGT 1021

      74 euLeuAspAlaCysGlyPheTyrTrpGlyProLeuSerValHisGlyAla 90
          |||:::   :::   :::||||||||||   :::|||   :::   :::
    1022 CCTTACAAAATGCTGAATGGTACTGGGGAGATATCTCGAGGGAAGAAGTG 1071

      91 HisGluArgLeuArgAlaGluProValGlyThrPheLeuValArgAsp.. 106
         :::|||:::||||||          ||||||||||||||||||||
    1072 AATGAAAAACTTCGAGATACAGCAGACGGGACCTTTTTGGTACGAGATGC 1121

     107 .SerArgGlnArgAsnCysPhePheAlaLeuSerValLysMetAlaSerG 123
          |||   :::   :::       ::::::|||:::::::::   :::::::
    1122 GTCTACTAAAATGCATGGTGATTATACTCTTACACTAAGGAAAGGGGGAA 1171

     123 lyProThrSerIleArgValHisPheGlnAlaGlyArgPheHisLeuAsp 139
          ::          |||:::::::                 ||||||   |||
    1172 ATAACAAATTAATCAAAATA.....................TTTCATCGAGAT 1203

     140 GlySerArgGlu.....................ThrPheAspCysLeuPheGluLe 151
         |||:::                         |||||||:::   :::   |||||
    1204 GGGAAATATGGCTTCTCTGACCCATTAACCTTCAGTTCTGTGGTTGAATT 1253

     151 uLeuGluHisTyrValAlaAlaProArgArgMetLeuGlyAlaProLeuA 168
          |:::::::||||||                  :::   :::      |||
    1254 AATAAACCACTACCGGAATGAATCTCTAGCTCAGTATAATCCCAAATTGG 1303

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      185 ..............AlaAlaValGlyArgGluAsnLeuAlaArgIlePr 196
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     1354 AAAGAAGATAATATTGAAGCTGTAGGGAAAAAA..................... 1386

      196 oLeuAsnProValLeuArgAspTyrLeuSerSerPhe 208
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 seq_name: /cgn1_6/ptodata/2/ina/5C_COMB.seq:US-08-539-005-1

 seq_documentation_block:
 ; Sequence 1, Application US/08539005
 ; Patent No. 5858686
 ;  GENERAL INFORMATION:
 ;    APPLICANT:  Schlessinger, Joseph
 ;    APPLICANT:  Skolnick, Edward Y.
 ;    APPLICANT:  Margolis, Benjamin L.
 ;    TITLE OF INVENTION:  NOVEL EXPRESSION CLONING METHOD FOR
 ;    TITLE OF INVENTION:  IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE
 ;    TITLE OF INVENTION:  KINASES AND NOVEL TARGET PROTEINS
 ;    NUMBER OF SEQUENCES:  50
 ;    CORRESPONDENCE ADDRESS:
 ;      ADDRESSEE:  PENNIE & EDMONDS
 ;      STREET:  1155 Avenue of the Americas
 ;      CITY:  New York
 ;      STATE:  New York
 ;      COUNTRY:  10036-2711
 ;      ZIP:  10036-2711
 ;    COMPUTER READABLE FORM:
 ;      MEDIUM TYPE:  Floppy disk
 ;      COMPUTER:  IBM PC compatible
 ;      OPERATING SYSTEM:  PC-DOS/MS-DOS
 ;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
 ;    CURRENT APPLICATION DATA:
 ;      APPLICATION NUMBER:  US/08/539,005
 ;      FILING DATE:  4-OCT-1995
 ;      CLASSIFICATION:  435
 ;    PRIOR APPLICATION DATA:
 ;      APPLICATION NUMBER:  US 08/167,035
 ;      FILING DATE:  16-DEC-1993
 ;      CLASSIFICATION:  435
 ;    ATTORNEY/AGENT INFORMATION:
 ;      NAME:  Coruzzi, Laura A.
 ;      REGISTRATION NUMBER:  30,742
 ;      REFERENCE/DOCKET NUMBER:  7683-062
 ;    TELECOMMUNICATION INFORMATION:
 ;      TELEPHONE:  (212) 790-9090
 ;      TELEFAX:  (212) 869-9741/8864
 ;      TELEX:  66141 PENNIE
 ;  INFORMATION FOR SEQ ID NO:  1:
 ;    SEQUENCE CHARACTERISTICS:
 ;      LENGTH:  3372 base pairs
 ;      TYPE:  nucleic acid
 ;      STRANDEDNESS:  double
 ;      TOPOLOGY:  unknown
 ;    MOLECULE TYPE:  cDNA
 ;    FEATURE:
 ;      NAME/KEY:  CDS
 ;      LOCATION:  43..2214
 US-08-539-005-1


 alignment_scores:
             Quality:  119.50                Length:    179
               Ratio:  1.258                   Gaps:      7
  Percent Similarity:  53.073    Percent Identity: 24.581

 alignment_block:
  US-08-962-560A-4 x US-08-539-005-1    ..

  Align seg 1/1  to: US-08-539-005-1  from: 1  to: 3372

       41 ArgProCysProAlaValProAlaProAlaProGlyAspThrHisPheAr 57
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    :::|||:::||||||:::|||         |||      |||
 946 CAGCCTGCACCAGCACTGCCTCCTAAACCACCAAAACCTACT..... 987

  57 gThrPheArgSerHisSerAspTyrArgArgIleThrArgThrSerAlaL 74
      |||      :::::::::             :::::::::
 988 .ACTGTAGCCAACAACGGT.............ATGAATAACAATATGT 1021

  74 euLeuAspAlaCysGlyPheTyrTrpGlyProLeuSerValHisGlyAla 90
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1022 CCTTACAAAATGCTGAATGGTACTGGGGAGATATCTCGAGGGAAGAAGTG 1071

  91 HisGluArgLeuArgAlaGluProValGlyThrPheLeuValArgAsp.. 106
     :::|||:::||||||            ||||||||||||||||||||
1072 AATGAAAAACTTCGAGATACAGCAGACGGGACCTTTTTGGTACGAGATGC 1121

 107 .SerArgGlnArgAsnCysPhePheAlaLeuSerValLysMetAlaSerG 123
      |||   :::   :::     :::::::|||:::::::::   :::::::
1122 GTCTACTAAAATGCATGGTGATTATACTCTTACACTAAGGAAAGGGGGAA 1171

 123 lyProThrSerIleArgValHisPheGlnAlaGlyArgPheHisLeuAsp 139
     ::        |||:::::::                  ||||||   |||
1172 ATAACAAATTAATCAAAATA.................TTTCATCGAGAT 1203

 140 GlySerArgGlu.............ThrPheAspCysLeuPheGluLe 151
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